

# Replacement Sheet

Figure 15A

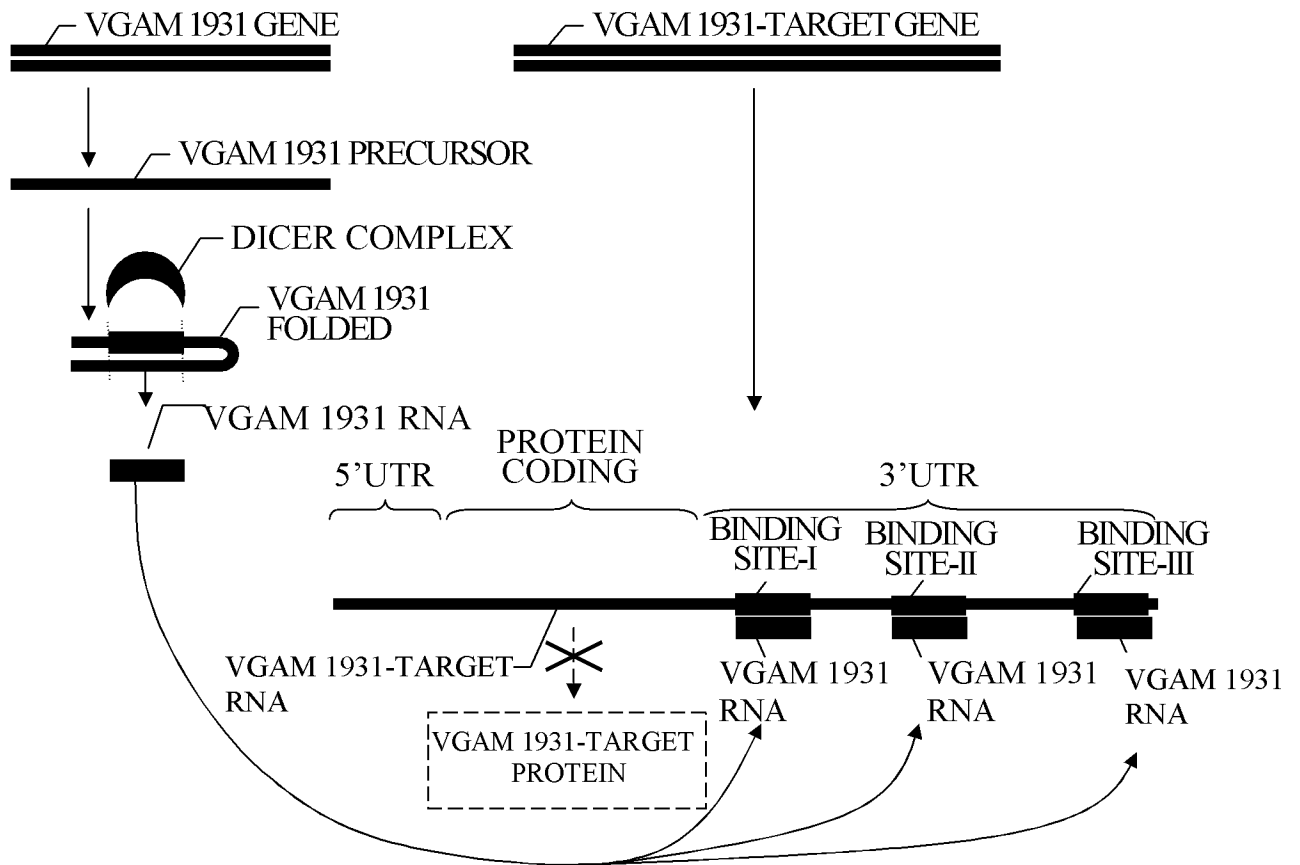


Figure 15B

CCCGTGGGGGGTCTTAGTGGAAGTGACGTGCTGTGAATA **SEQ ID:1917**  
CAGGTCCATAGCACCGCTATCCACTATGTCTCGCCCGGG

TAGCACCGCTATCCACTATGTCTC

**SEQ ID: 4642**

Figure 15C

T	G	TCT	-	AC	A	A
CCCG	GG	GGG	TAGTGGA	AGTG	GTGCTGTG	AT C
GGGC	CC	CTC	ATCACCT	TCGC	CACGATAC	TG A
-	G	TGT	A	--	C	G

# Replacement Sheet

Figure 15D/1

COL6A1 BINDING SITE1	5' TAGCACCG ATC GTCT 3'	SEQ ID:4642
	3' ATCGTGGC TAG CAGA 5'	SEQ ID:7584
	CT- CACTAT               AAT AGCTCC	
SFRS1 BINDING SITE1	5' GCA GCTATCCACT GTC 3'	SEQ ID:4642
	3' CGT CGATGGGTGG CAG 5'	SEQ ID:13801
	CC AT-               AA ATT	
HIP12 BINDING SITE1	5' TAGC CCG CC CTATGTCTC 3'	SEQ ID:4642
	3' GTCG GGC GG GATACAGAG 5'	SEQ ID:32922
	A- CTAT A                 CG CC-- C	
ZNF212 BINDING SITE1	5' TAGCACCG TCC TG TCTC 3'	SEQ ID:4642
	3' ATCGTGGC AGG AC AGAG 5'	SEQ ID:14557
	CTA ACTA -                  AC- GG-- G	
FLJ20436 BINDING SITE1	5' GC CCG TATCCACT TGTCT 3'	SEQ ID:4642
	3' CG GGT ATAGGTGA ACAGA 5'	SEQ ID:19472
	A C-- A                  - TTT C	
KIAA1622 BINDING SITE1	5' TAGCA C GCT TC TATGTCTC 3'	SEQ ID:4642
	3' ATTGT G CGA AG ATACAGAG 5'	SEQ ID:27766
	C - A CAC                     A T C TA-	
LOC51312 BINDING SITE1	5' GC GCTATCCA TAT TC 3'	SEQ ID:4642
	3' CG CGATAGGT ATA AG 5'	SEQ ID:20659
	ACC C G                 A-- - A	
LOC57105 BINDING SITE1	5' TAGC CG TA CTATGTCTC 3'	SEQ ID:4642
	3' ATCG GC AT GATATAGAG 5'	SEQ ID:21639
	AC C TCCA                  CA A CAAC	
LOC146603 BINDING SITE1	5' GCA GCT TCCACTATGT 3'	SEQ ID:4642
	3' TGT CGA AGGTGATACA 5'	SEQ ID:38215
	CC A               TA C	

# Replacement Sheet

Figure 15D/2

		G	ATC	-	A		
	5'	ACC	CT		CACT	TGTCTC	3'
LOC145761							SEQ ID:4642
BINDING SITE1	3'	TGG	GA		GTGA	ACAGAG	5'
		G	AAA		C		SEQ ID:40584
			--		CT	CAC	
	5'	TAGCA	CCG		ATC	TATGTCTC	3'
LOC202986							SEQ ID:4642
BINDING SITE1	3'	ATCGT	GGT		TAG	ATACAGAG	5'
		TA	T-		CA-		SEQ ID:43470